Replacement paragraph for p.15, lines 8-11:

Fig. 1 presents DNA sequence data (SEQ ID NO: 2) generated using M13mp18 containing a 115 bp SauAI fragment from lambda inserted a the BamHI site and Cy5.5 ddGTP, ddATP, ddTTP, and ddCTP dye terminators.

This listing of claims will replace all prior versions, and listings, of claims in the application:

## Listing of Claims:

1. (Original) A kit for DNA sequencing comprising:

a first, second, third and fourth dye terminator molecule, each of the dye terminator molecules comprising a dye molecule, a linker of at least 10 atoms in length and either ddATP, ddCTP, ddGTP or ddTTP as a mono or tri-phosphate and a thermostable DNA polymerase.

- 2. (Original) The kit of claim 1, wherein said polymerase is a thermostable DNA polymerase that has an altered dNMP binding site so as to improve the incorporation of dideoxynucleotides relative to the natural polymerase.
  - 3. (Original) A compound of formula (I)

wherein A is a cyanine dye of the structure

$$\begin{array}{c|c}
R_3 & X & H & R_7 \\
 & & C = C & C & R_2
\end{array}$$

$$\begin{array}{c|c}
R_4 & & & \\
R_5 & & & \\
R_1 & & & & \\
\end{array}$$

$$\begin{array}{c|c}
R_7 & & H & & \\
R_7 & & & & \\
R_1 & & & & \\
\end{array}$$

and the curved lines represent carbon atoms necessary for the formulation of cyanine dyes, X and Y are selected from the group consisting of O, S, and CH<sub>3</sub>-C-CH<sub>3</sub>, m is an integer selected from the group consisting of 1, 2, 3, and 4; R1, R2, R3, R4, R5, R6 and R7 are independently selected from the group consisting of H, OH, CO<sub>2</sub>H, sulfonic acid or sulfonate groups, esters, amides, ethers, alkyl or aryl groups and B, and one R1, R2, R3, R4, R5, R6 or R7 is B;

B is a linker of at least 10 atoms in length wherein the atoms are selected from the group consisting of carbon, nitrogen, oxygen, substituted carbon, and sulfur and the linker is attached at one end to A and at the other end to C;

C is a dideoxynucleotide selected from the group consisting of

and wherein said linker is covalently bonded to said dideoxynucleotide at position 7 for ddA and ddG and at position 5 for ddC and ddT and wherein r is a mono or tri-phosphate.

- 4. (Original) The compound of claims 3, wherein said linker is selected from the group consisting of
  - -C≡C- CH<sub>2</sub>-NH-CO- (CH<sub>2</sub>)  $_5$ -NH-CO-,
  - -C≡C- CH<sub>2</sub>-NH-CO- (CH<sub>2</sub>) 9-NH-SO<sub>2</sub>-,
  - -C $\equiv$ C- CH<sub>2</sub>-NH-CO- (CH<sub>2</sub>) <sub>10</sub>-NH-CO-,
  - -C≡C- CH<sub>2</sub>-NH-CO- (CH<sub>2</sub>) 5-,
  - -C≡C- CH<sub>2</sub>-NH-CO- (CH<sub>2</sub>) 5 -NH-CO- (CH<sub>2</sub>) 5-, and
  - -C $\equiv$ C- CH<sub>2</sub>-NH-CO- (CH<sub>2</sub>) 5 -NH-CO- (CH<sub>2</sub>) 10 -NH-CO- .

## 5. (Original) A compound of the formula (II):

## 6. (Original) A compound of the formula (III):

## 7. (Original) A compound of the formula (IV):

8. (Original) A compound of the formula (V):

- 9. (Orignal) A deoxyribonucleic acid sequence containing the compound of formula I.
- 10. (Previously amended) A deoxyribonucleotide sequence containing a compound of formula II, III, IV, or V.

- 11. (Original) A kit for DNA sequencing comprising compounds of formula II, III, IV, and V.
  - 12. (Original) The kit of claim 11, further comprising a thermostable DNA polymerase.
- 13. (Original) The kit of claim 12, wherein said polymerase is a thermostable DNA polymerase that has an altered dNMP binding site so as to improve the incorporation of dideoxynucleotides relative to the natural polymerase.
  - 14-17. Cancelled.
  - 18. (New) The kit of claim 1, wherein said linker is a linker of 10 to 25 atoms.
- 19. (New) The kit of claim 18, wherein the dye molecule on at least one or said dye terminator molecules is a cyanine dye.
- 20. (New) The kit of claim 18, wherein the dye molecules on each of said dye terminator molecules is a cyanine dye.
  - 21. (New) The compound of claim 3, wherein said linker is a linker of 10 to 25 atoms.